CLASS SUBCLASS APPROVED O.G. FIG. DRAFTSMAN à

GAATTCAACT TCTCCATACT TTGGATAAGG AAATACAGAC ATGAAAATC TCATTGCTGA GTTGTTATTT AAGCTTGCCC AAAAAGAAGA AGAGTCGAAT CTTAAGTIGA AGAGGIATGA AACCIATICC ITTAIGICIG IACTITITAG AGIAACGACI CAACAAIAAA ITCGAACGGG ITITICITCI ICTCAGCTIA

CTIGACACAC GCGTCCATCT TCGAAACCTC TAATAGCAGT GACGTTACGA AGCGTTATAC CGCGTTTTAC TGGTTGTCGC CAACTAACTA GTCCATCTCC TCGCAATATG GCGCAAAATG ACCAACAGCG GTTGATTGAT CAGGTAGAGG GAACTGTGTG CGCAGGTAGA AGCTTTGGAG ATTATCGTCA CTGCAATGCT 101

GCATTCCTGA CGACGATACG GAGCTGCTGC GCGATTACGT AAAGAAGTTA TTGAAGCATC CTCGTCAGTA CCCGCGACAT GCTCCATTTC GGGCTACGGT CGTAAGGACT GCTGCTATGC CTCGACGACG CGCTAATGCA TTTCTTCAAT AACTTCGTAG GAGCAGTCAT GGGCGCTGTA CGAGGTAAAG CCCGATGCCA 201

GTACGCAAGT TTTTCAATTA GAAAAGTTGT CGACAGTATT TCAACAGTGC CGGCTCTGAA TATCAGCGAA ACAAAATAA AAAATTACAT AAACATTGAT CATGCGTTCA AAAAGTTAAT CTTTTCAACA GCTGTCATAA AGTTGTCACG GCCGAGACTT ATAGTCGCTT TGTTTTTATT TTTTAATGTA TTTGTAACTA 301

401 TCACGTAAAA AGGGTATCTA GAGGTTGAGG TGATTTT AGTGCATTTT TCCCATAGAT CTCCAACTCC ACTAAAA STII xbaI Trp SD

AAA AGA Phe Ser ATG AAA AAG AAT ATC GCA TTT CTT GCA TCT ATG TTC GTT TAC TTT TTC TTA TAG CGT AAA GAA GAA CGT AGA TAC AAG CAA Met Lys Lys Asn Ile Ala Phe Leu Leu Ala Ser Met Phe Val

486 ATT GCT ACA AAT GCC TAT GCA (SEQ ID NO: 13)

TGT TTA CGG ATA CGT TAA CGA

Ala Thr Asn Ala Tyr Ala (SEQ ID NO: 14) 17 Ile

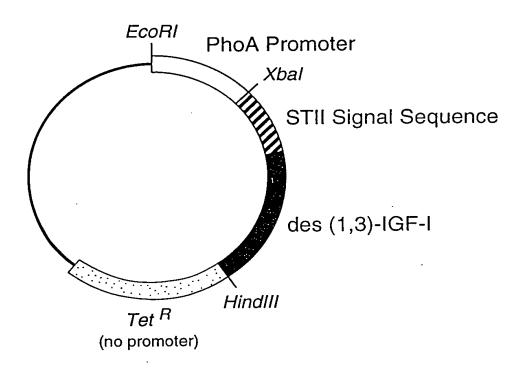
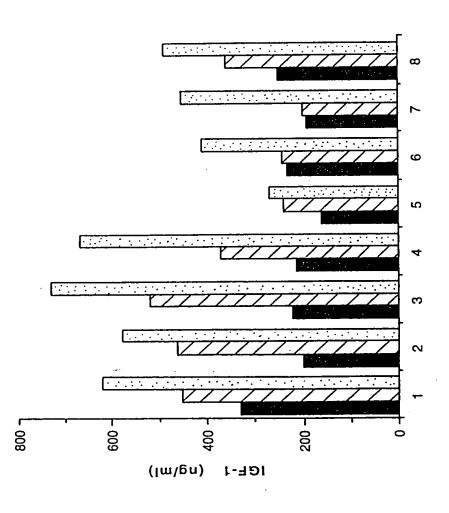
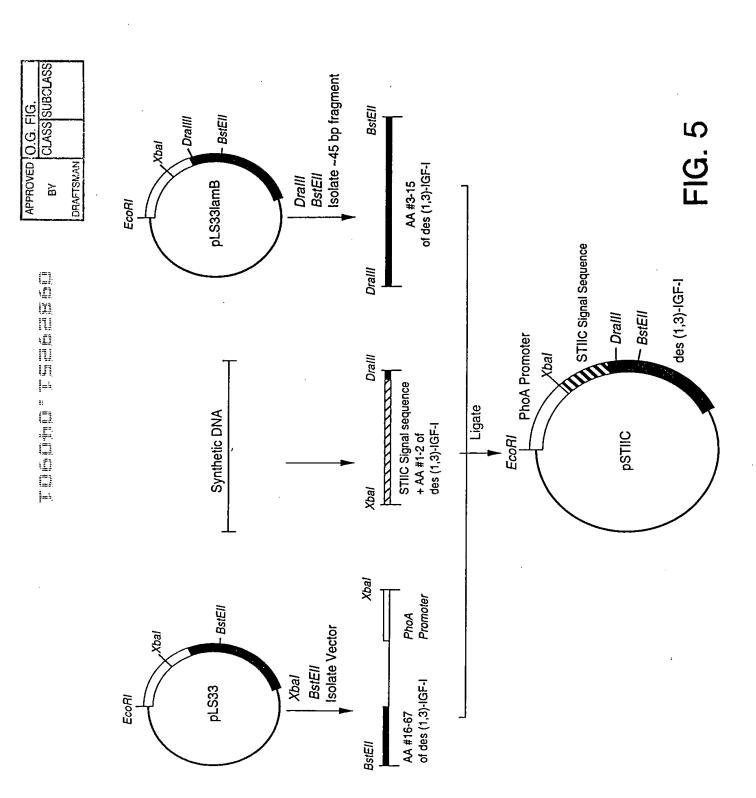
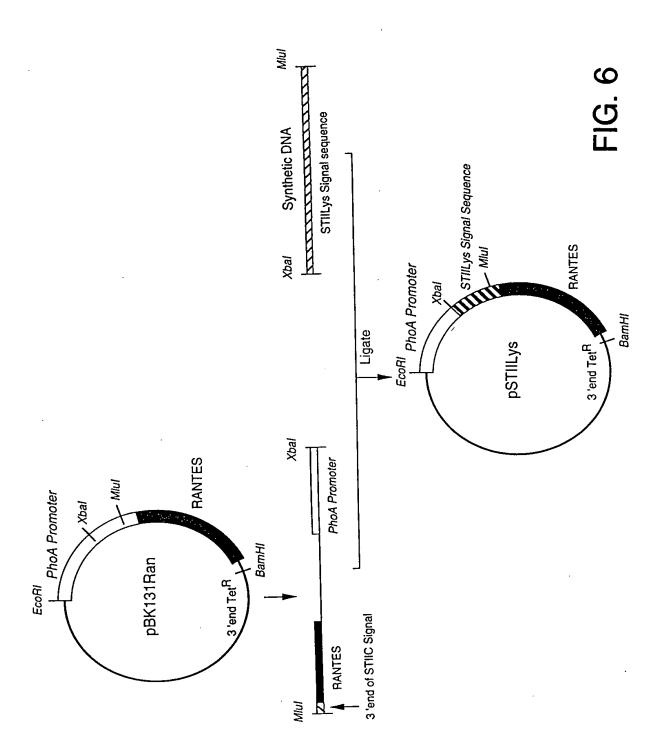


FIG. 2



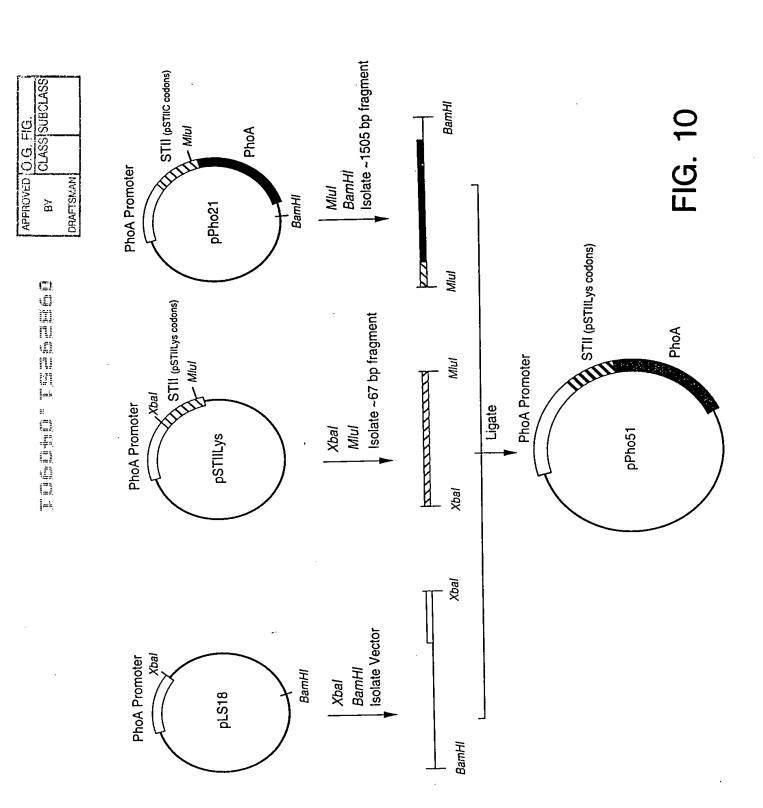
Experiment





APPROVED O.G. FIG. BY CLASS SUBCLASS DRAFTSMAN	PhoA Promoter STII (pSTiIC codons) pPho21 PhoA	Miul BamHI Isolate ~1505 bp fragment	BamHI .	FIG. 8
The state of the s	EcoRI PhoA Promoter STII STII MIUI	EcoRI Miul Isolate ~496 bp fragment	EcoRI Miul Miul Miul Ligate PhoA Promoter	pPho31 PhoA
	EcoRI pBR322.	EcoRI BamHI Isolate Vector	BamHI EcoRI	

APPROVED O.G. FIG. BY CLASS SUBCLASS DRAFTSMAN	PhoA Promoter STII (pSTIIC codons) pPho21 PhoA	Miul BamHl Isolate ~1505 bp fragment	BamHI odons)	FIG. 9
read miner games, county grows, county presser,	EcoRI PhoA Promoter STII (pSTIIBK #131 codons) pNGF131 NGF	EcoRI MIU! Isolate ~484 bp fragment	EcoRI Miul Miul Miul PhoA Promoter PhoA Promoter STII (pSTIIBK #131codons)	pPho41
	EcoRI pBR322	EcoRI BamHI Isolate Vector	BamHI EcoRI	



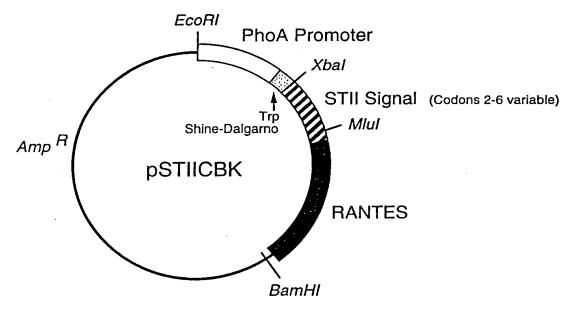
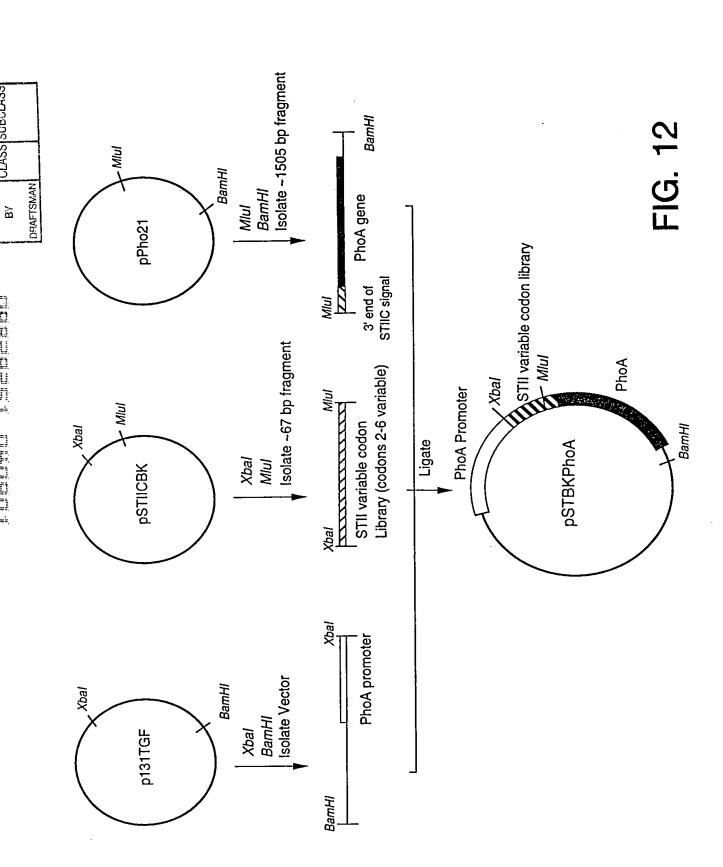


FIG. 11



griff genis genis consi genis consi grino. Al genis del Berlin del

APPROVED O.G. FIG.

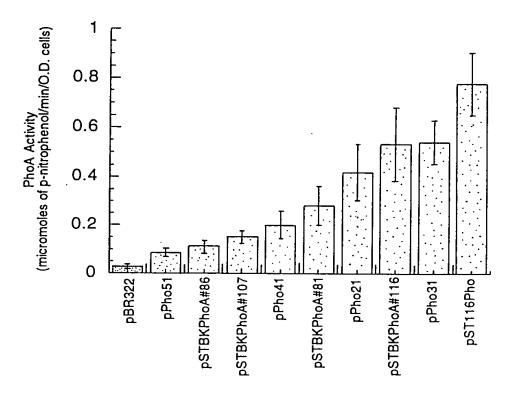


FIG. 13

0.G. FIG.	CLASS SUBCLASS	
APPROVED	BY	DRAFTSMAN

The final track from the first track from the first

pPho41 (STIIBK#131) TCTAGAATT ATG AAG AAT ATT GCG TTC CTA CTT GCC TCT ÀTG TTT GTC pSTBKPhoA#107 TCTAGAATT ATG AAG AAA AAC ATC GCT TTT CTT GCA TCT ATG TTC GTT pPho31 (Wild type STII + Mlul site) TCTAGAGGTIGAGGTGATTTT ATG AAA AAG AAT ATC GCA TTT CTT CTT GCA TCT ATG TTC GTT pPho51 (STIILys - unless otherwise noted this sequence is the TIR=1 used in the examples)

Sequence is the TIR=1 used in the examples)

TCTAGAATT ATG AAG AAT ATC GCA TTT CTT CTT GCA TCT ATG TTC GTT pSTBKPhoA#81 TCTAGAATT ATG AAA AAA AAC ATT GCC TTT CTT GCA TCT ATG TTC GTT pPho21 (STIIC) TCTAGAATT ATG AAA AAG AAT ATC GCA TTT CTT GCA TCT ÄTG TTC GTT TCTAGAATT ATG AAA AAA AAC ATC GCA TTT CTT CTT GCA TCT ATG TTC GTT pSTBKPhoA#116_

FIG. 14A

dST116Pho Tctagaggttgaggtgatttt atg aaa aaa aac atc gca ttt ctt gca tct ätg ttc gtt

TCTAGAATT ATG AAA AAG AAC ATA GCG TTT CTT CTT GCA TCT ATG TTC STT

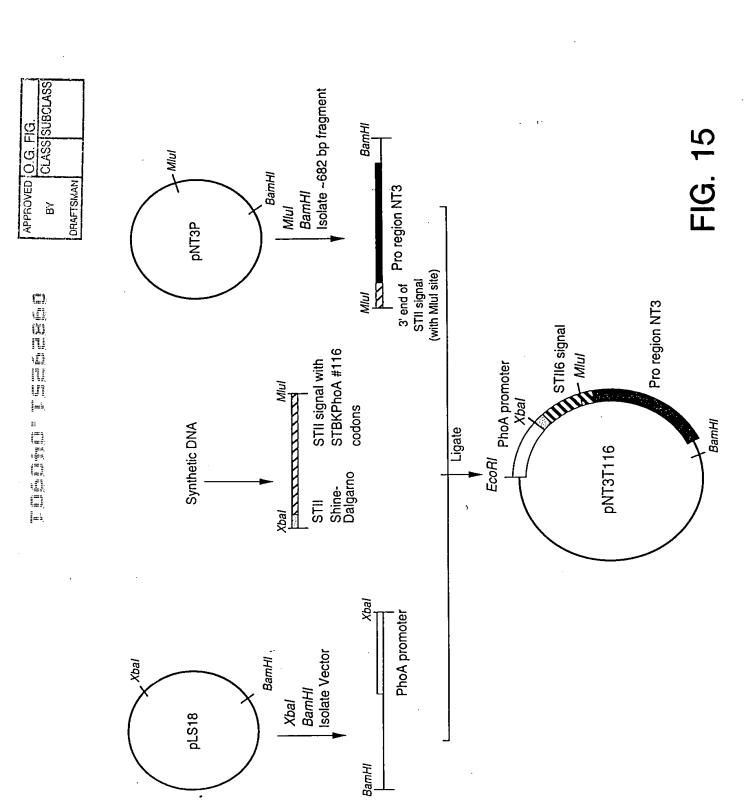
pSTBKPhoA#86

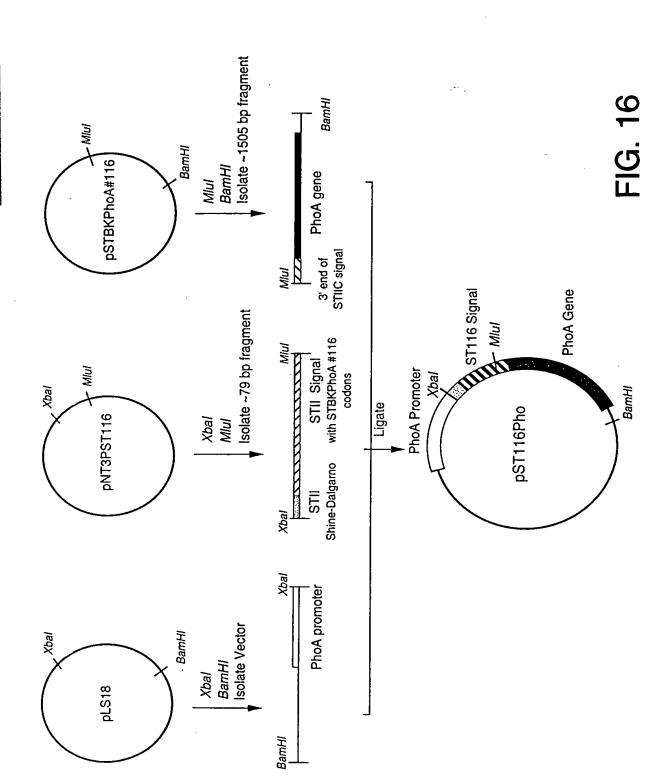
APPROVED O.G. FIG. BY CLASS SUI	O.G. FIG.	OLASS SUBCLASS	
	APPROVED	λ.	DRAFTSMAN

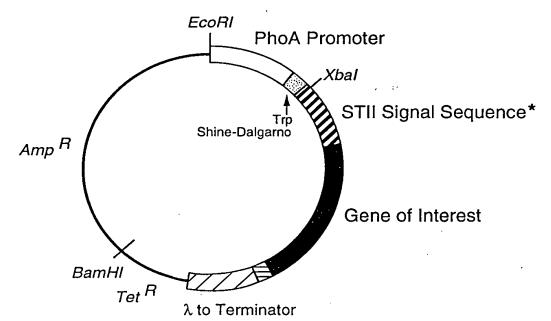
TIR RELATIVE STRENGTH	7	8	1	6	77	2	1	13
(SEQ ID NO:15)	(SEQ ID NO:16)	(SEQ ID NO:17)	(SEQ ID NO:18)	(SEQ ID NO:19)	(SEQ ID NO:20)	(SEQ ID NO:21)	(SEQ ID NO:22)	(SEQ ID NO:23)
TIT TCT ATT GCT ACA AAY GCS TAT GCM*	TTT TCT ATT GCT ACA AAC GCG TAT GCM	TTT TCT ATA GCT ACA AAC GCG TAT GCM	TTT TCT ATT GCT ACA AAC GCG TAT GCM	TTT TCT ATT GCT ACA AAC GCG TAT GCM	TTT TCT ATT GCT ACA AAC GCG TAT GCM	TTT TCT ATT GCT ACA AAC GCG TAT GCM	TTT TCT ATT GCT ACA AAC GCG TAT GCM	TTT TCT ATT GCT ACA AAC GCG TAT GCM

may vary in The codons for the last four amino acids of this sequence may distributed by the example, in the IGF-1, RANTES secretion plasmids, the sequence is AAT GCC TAT GCA. The last codon for the last amino acid in every sequence listed the examples of protein secretion - GCC and GCA were both used.

FIG. 14B

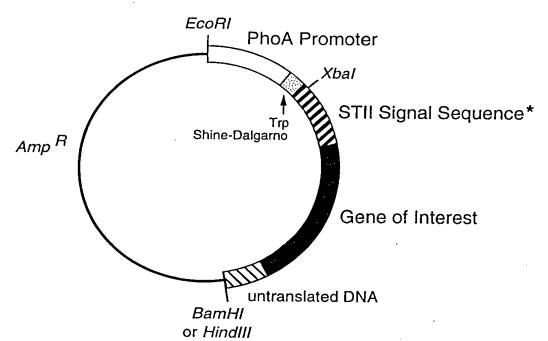






* One of the nucleotide sequences listed in Figure 14 (STII Shine-Dalgarno may also be included).

FIG. 17



* One of the nucleotide sequences listed in Figure 14 (STII Shine-Dalgarno may also be included).

FIG. 18